

#4

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 (T)
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Fig. 1

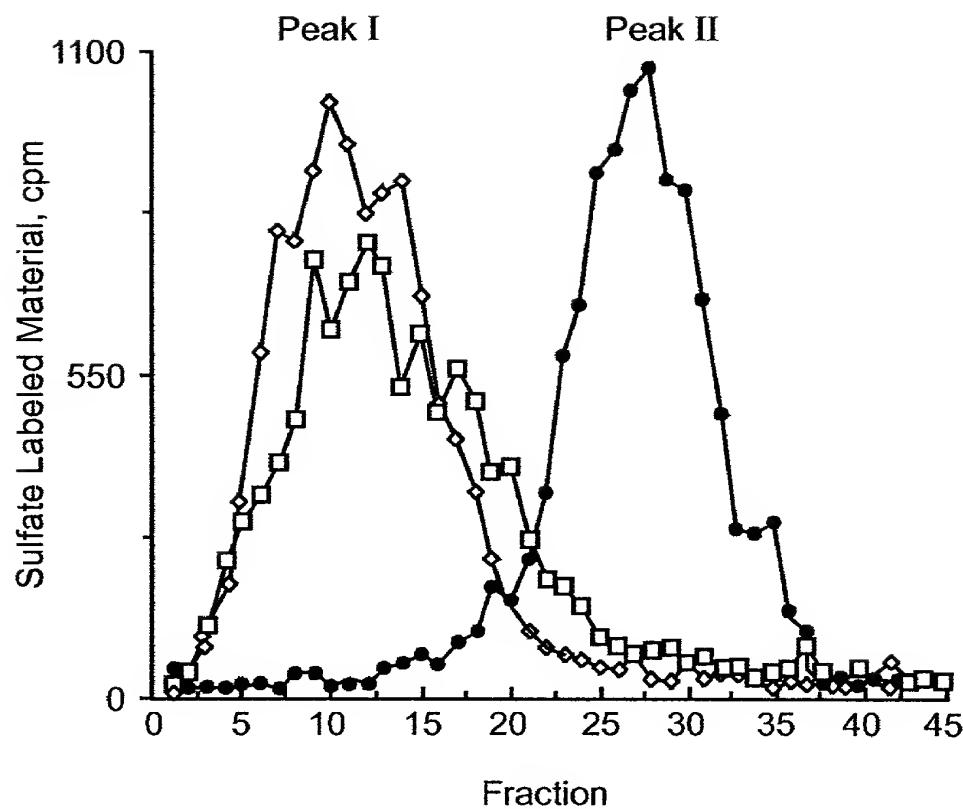


Fig. 2

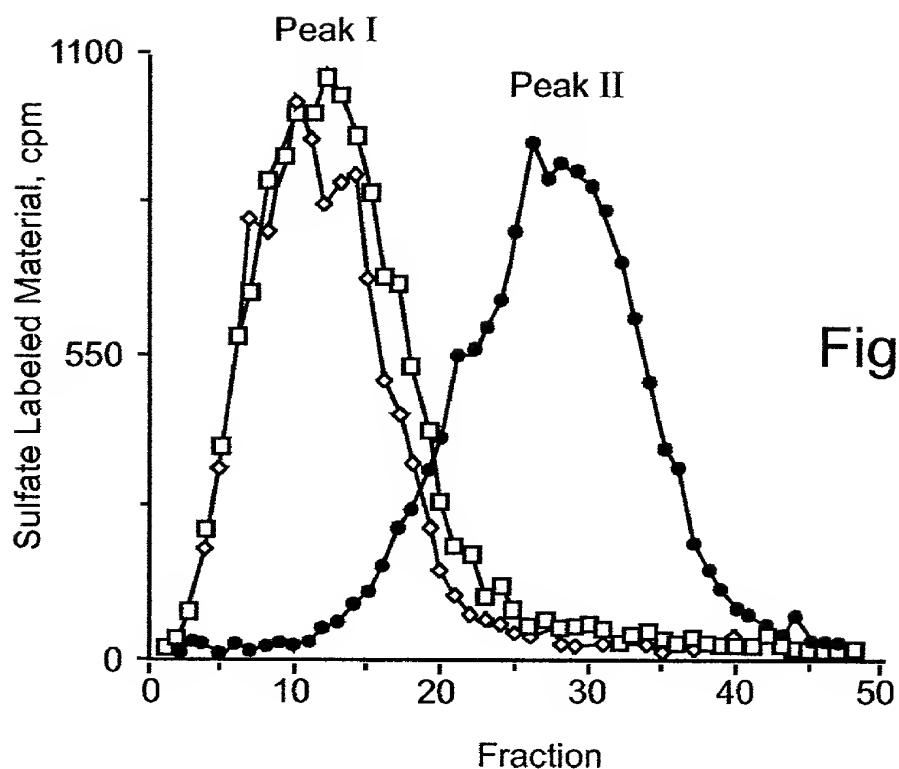


Fig. 3a

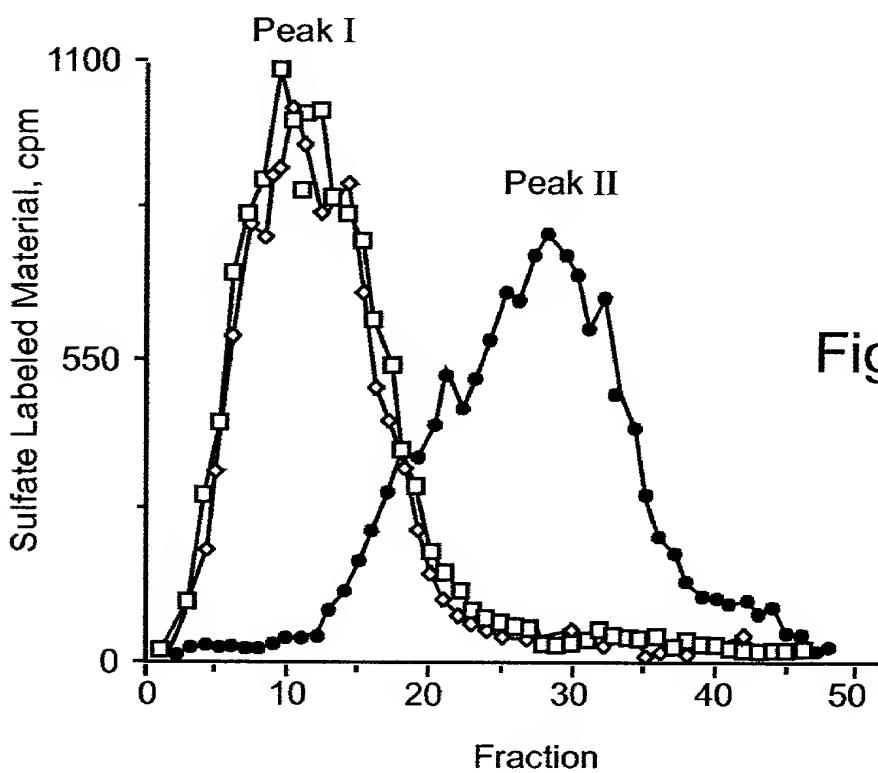
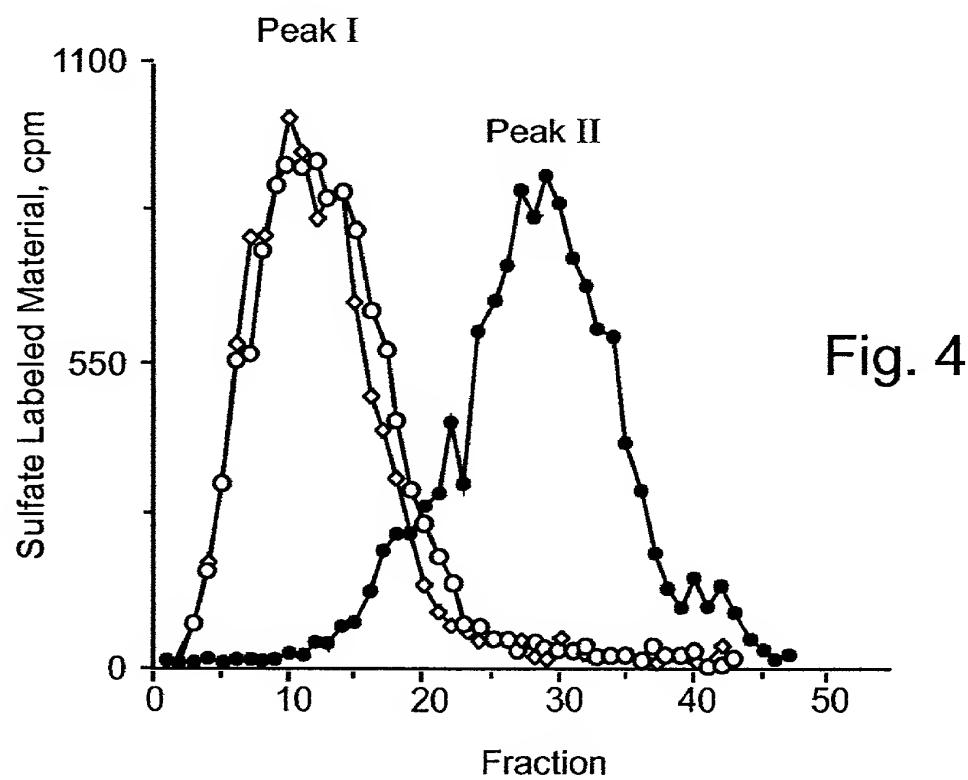


Fig. 3b



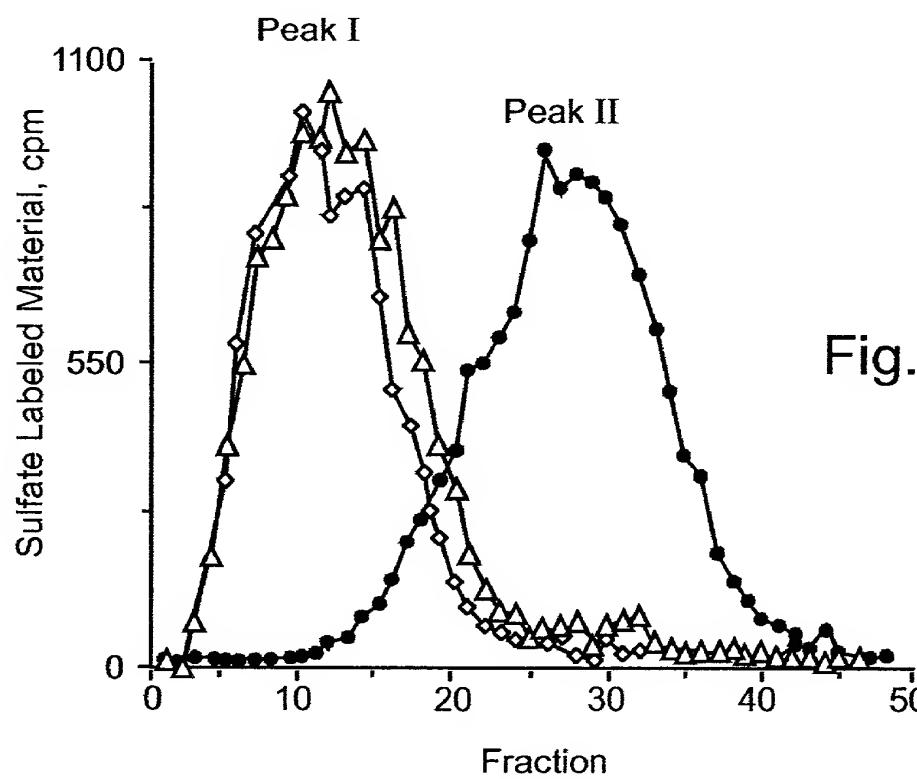


Fig. 5a

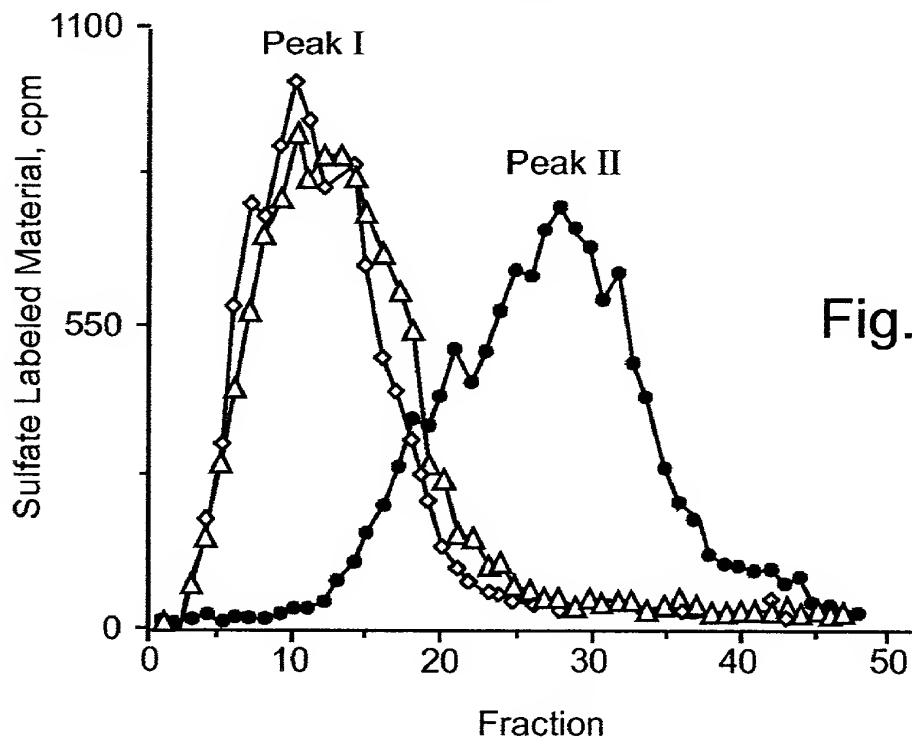
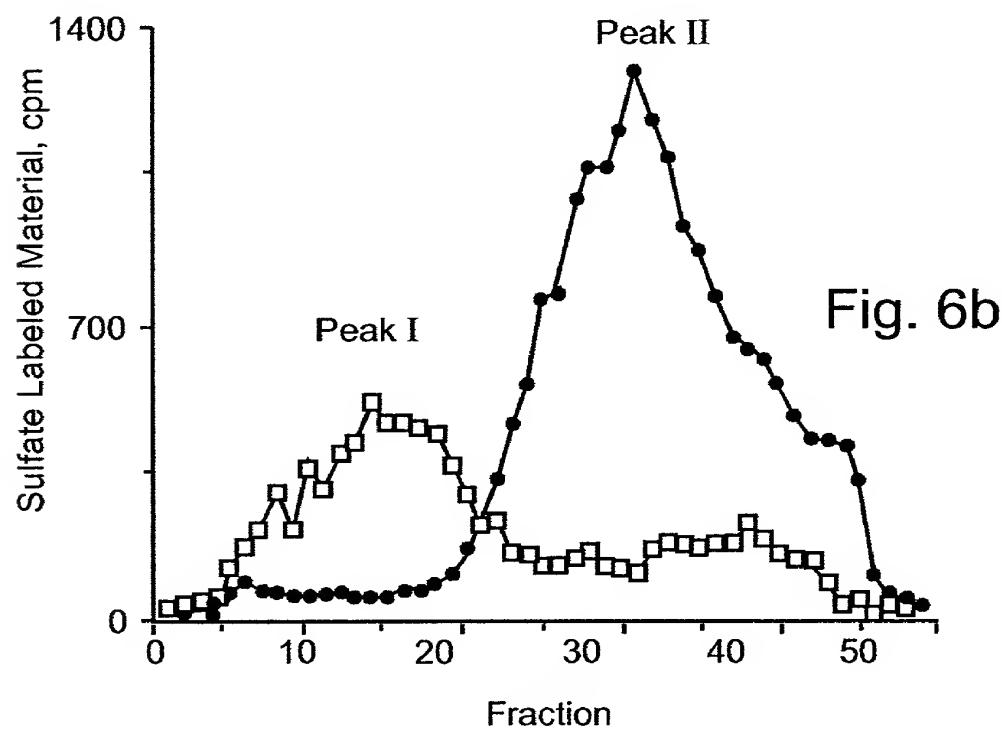
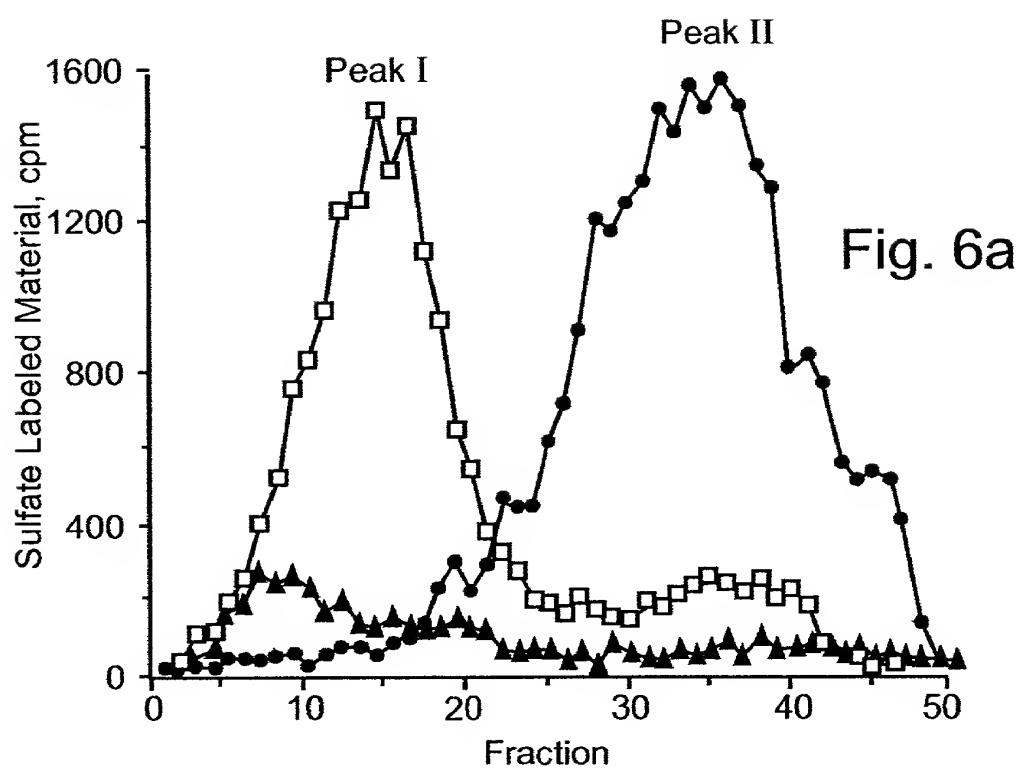


Fig. 5b



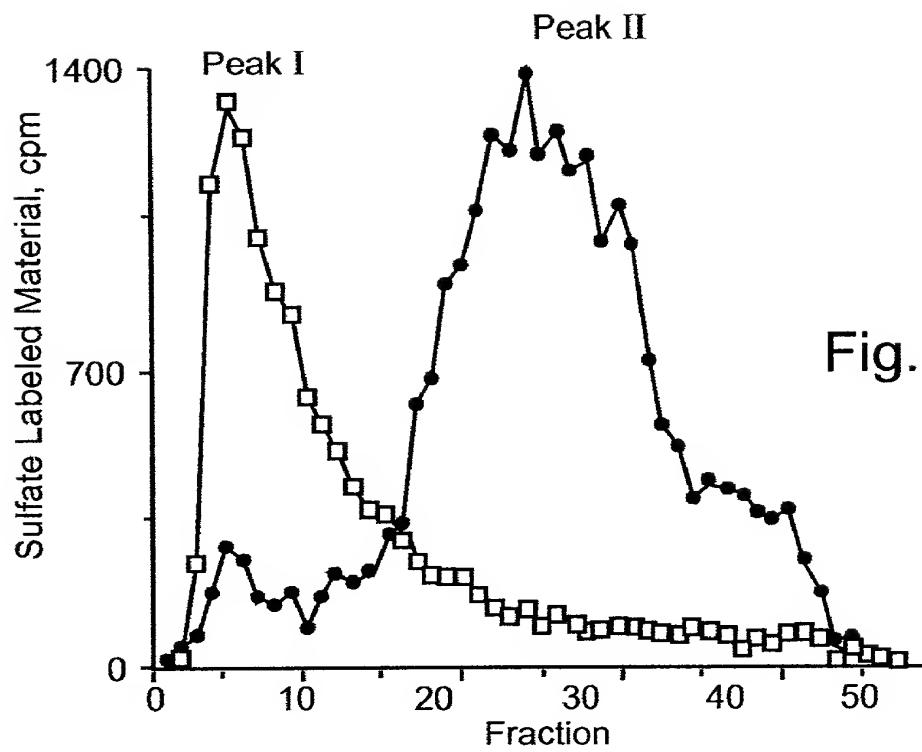


Fig. 7a

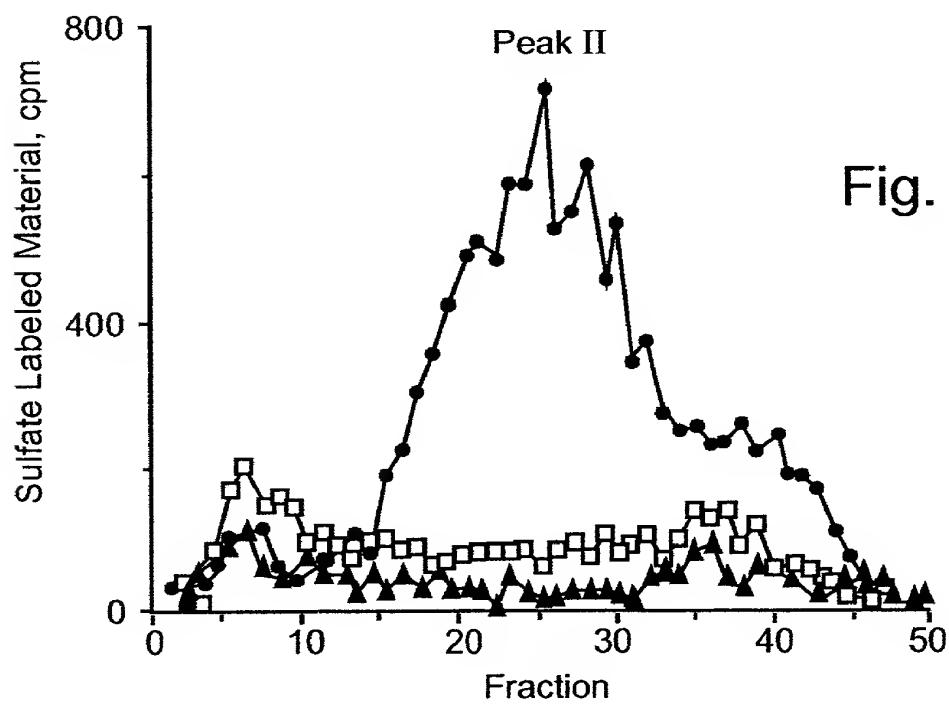


Fig. 7b

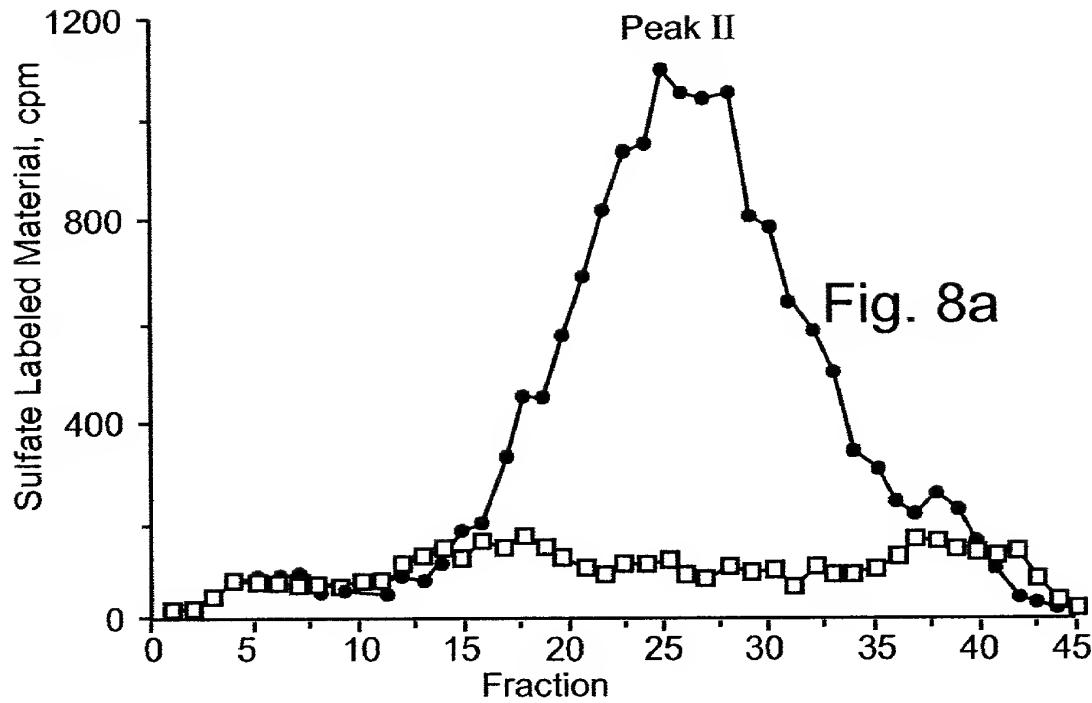


Fig. 8a

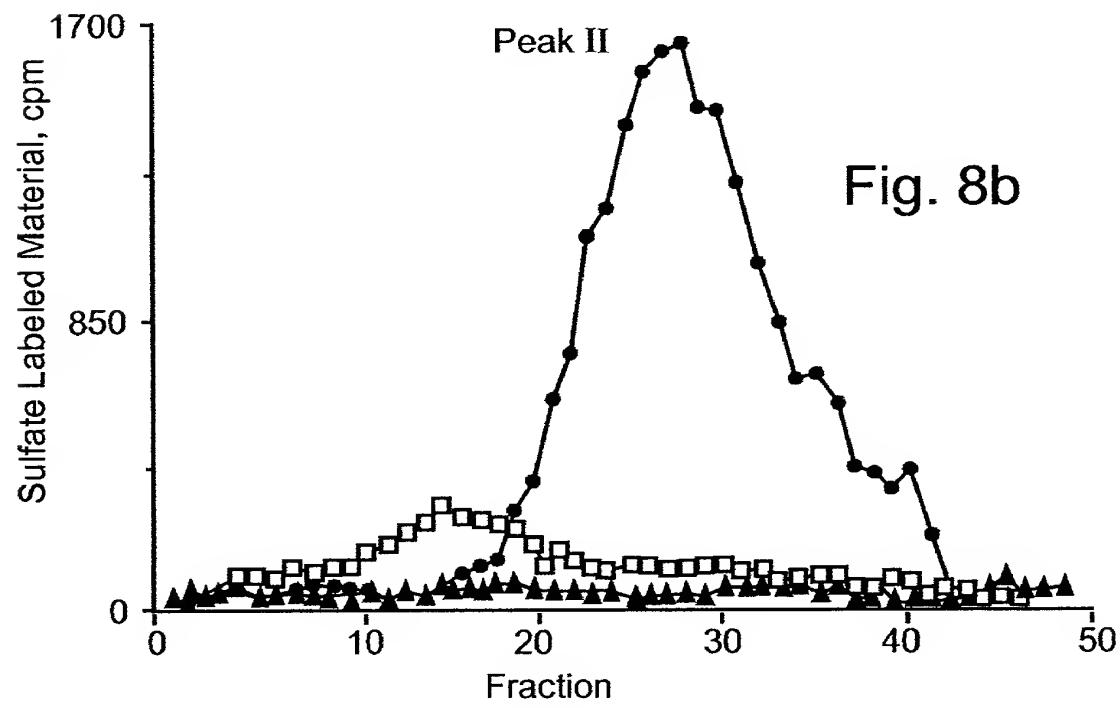
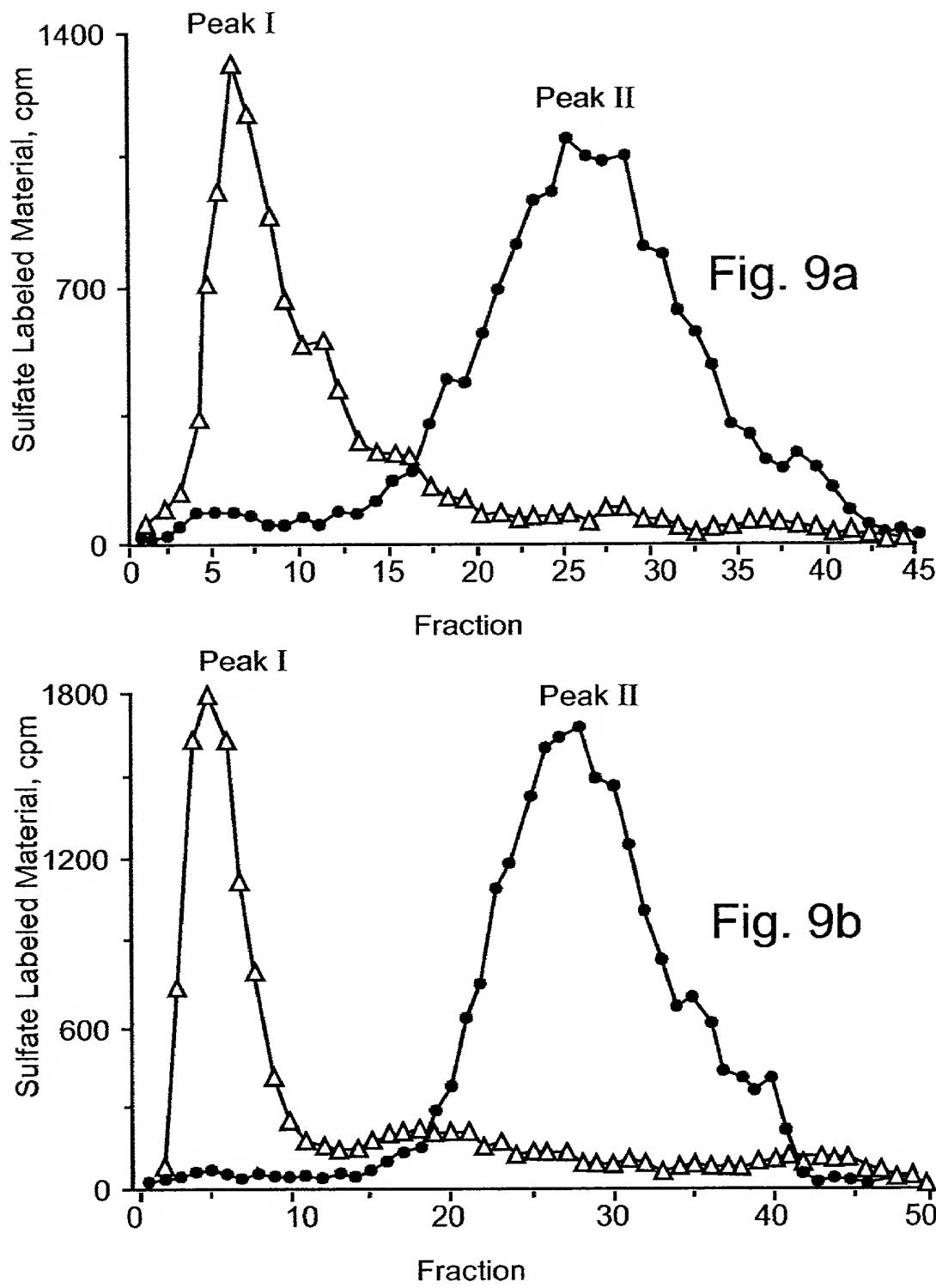


Fig. 8b



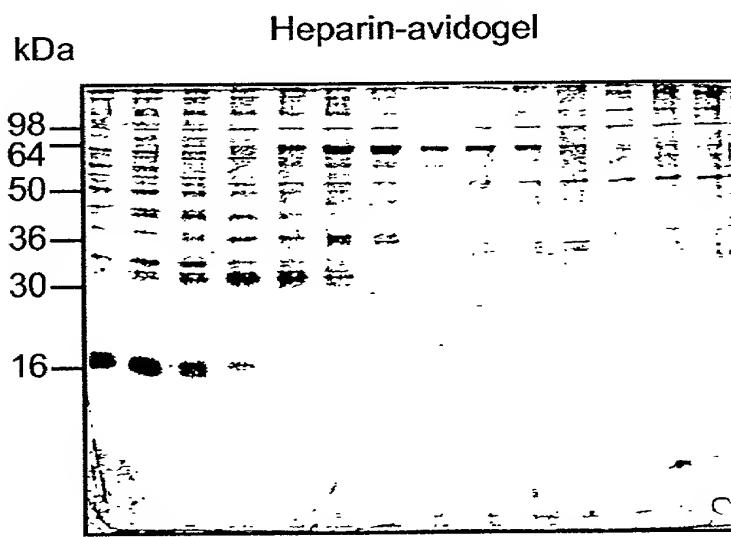
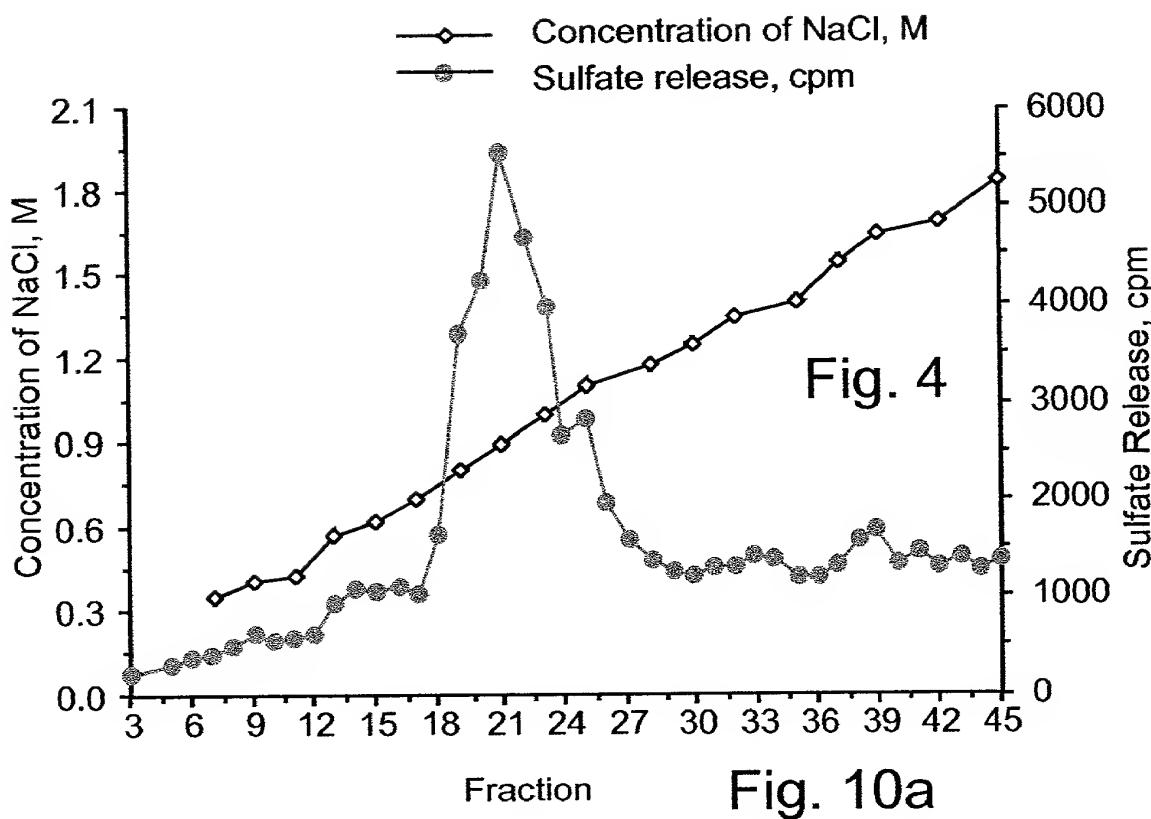


Fig. 10b

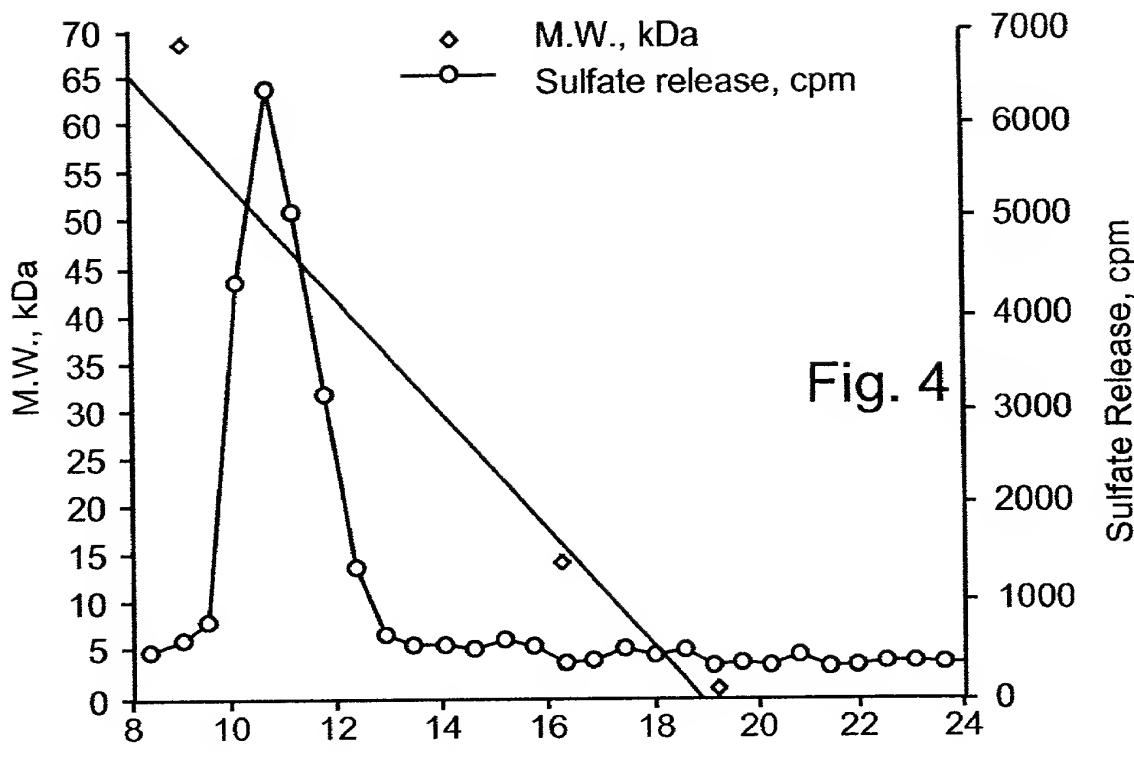


Fig. 4

Fig. 11a

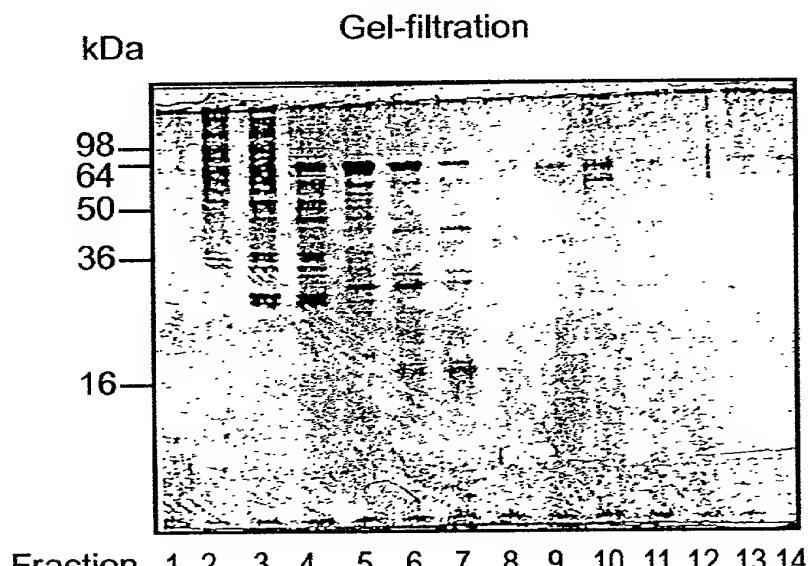


Fig. 11b

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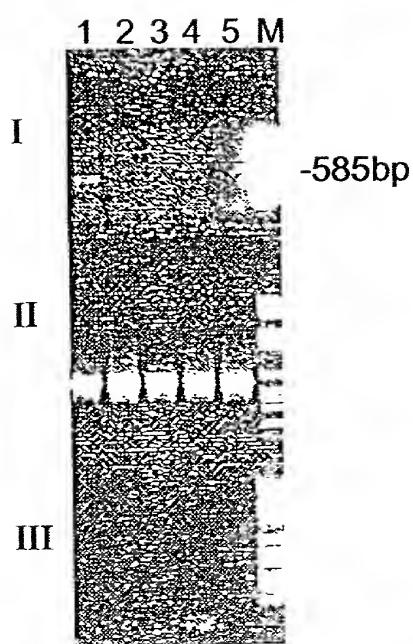


Fig. 12a

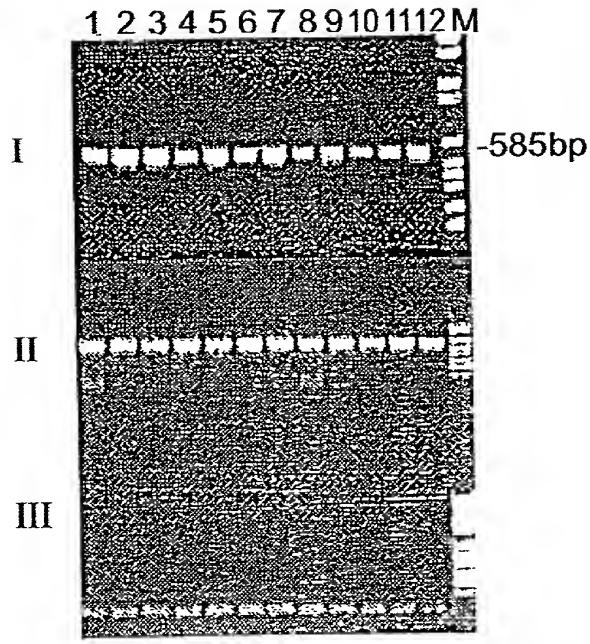


Fig. 12b

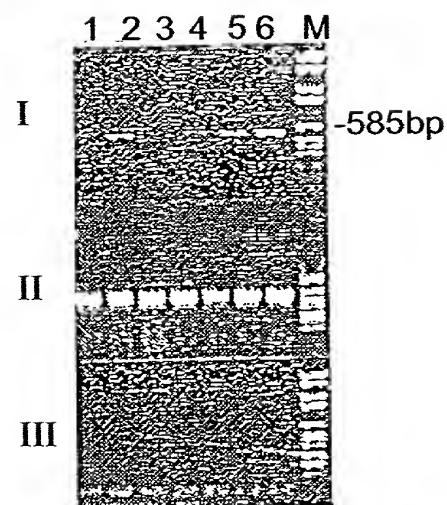


Fig. 12c

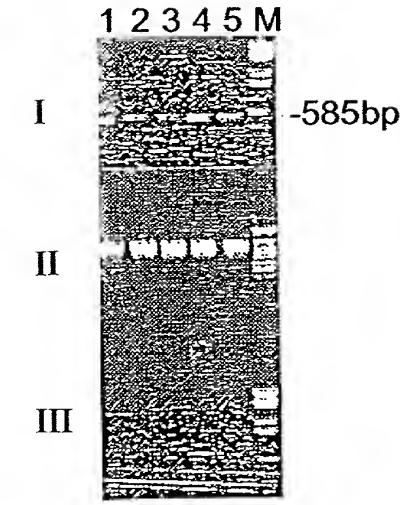


Fig. 12d

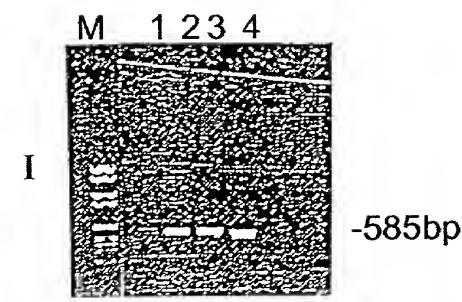


Fig. 12e

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mouse	GCACCCCTTGCTGTCCAACACACCTTGCAGCTGGCTTATGTGGCTGGATAA	100
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human	GTTAATGGCAAGCGTGCAAGGTTCAAAGAGAAGGAAGCTCGAGTATACC	1365
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human	CTGTATGCCATAAACCTCCATAACGTCAACAGTACTTGCACCTTACCTA	1465
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mouse	CGGATGGATTACTTCCAAATCTGTCCAACGTGAACGGTCAAATTCTGAAG	500
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Fig. 13

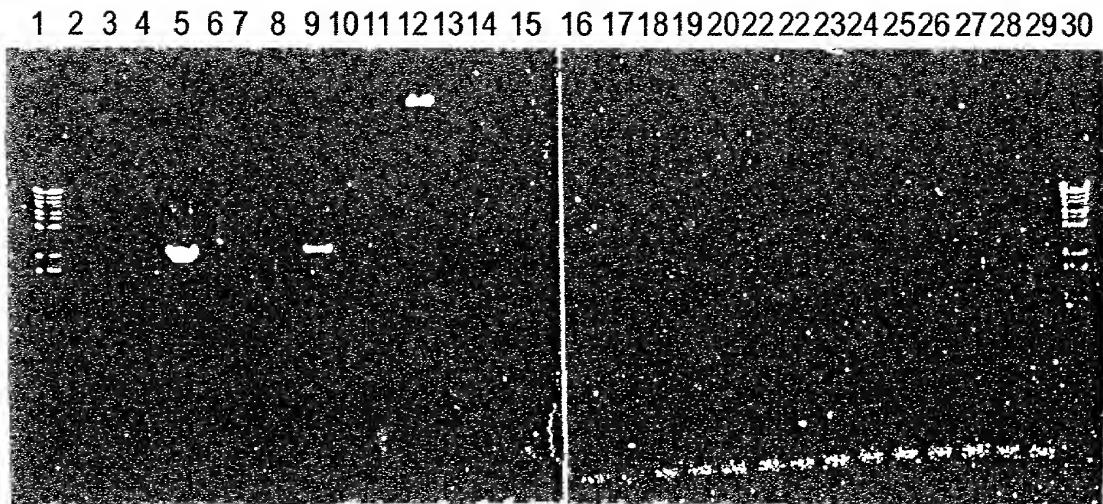


Fig. 14

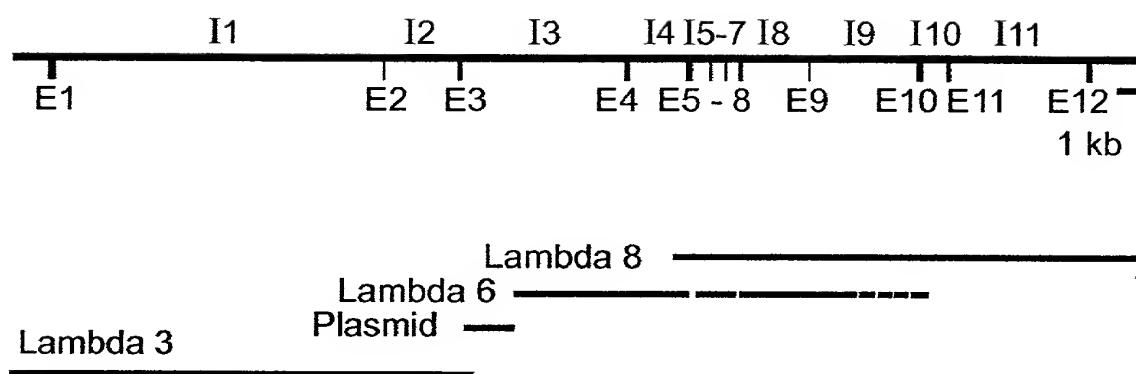


Fig. 15

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Fig. 16

Fig. 16
(continued)

Fig. 16
(continued)

Fig. 16
(continued)

Fig. 16
(continued)

Fig. 16
(continued)

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Fig. 16
(continued)

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gtaatgtactgccaccaaacggctgcgtatattggcaagacttacc
ttatttgaatctcaagtttctcttagaaaaatgagggtggaggtaagca
taggctgtatgtctaaagcctccatactgcctaaactgtggctctaag
atccagtagaatgtgggtcacaggactctaggagctttcaaaaa
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cagcaaatcaaaatacctaagcactgtcaagagtgttctgccta
gtggctttatagttaatattaaatagttaattttttttttttggagac
agagtctgtctgttaccaggctgcaggcactggcacaatctcggt
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agtagctggactacaggctgcactgcacccagactaattttgtat
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aggcatgagccactgcacccagcttaatagctaatattatatttc
tatagttattcaagtaattcaggccaaagacttagaaaacaaaaa
ccacttttaaggagaaaagggtgttagttgcacatagatagatctt
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atagatatacatgaaaatttggaggacttattatgtcatatgataatcaat
ttaaagacaacactaaaattatattgttgcactctcaaaaagtt
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aataggcataatgatgtccatagatgtttagtttttccct
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ccaagtagctgtggactacaggctgcattaccatcatgcccggtaatt
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tgaactcctggctcaagtgtatccggccgcctcggcctccaaagtgc
ggatgacaggcatgagccactgcacccctggcgaagtattt
aatggttacataggacataactaaacattttatgtctatataatgaagt

Fig. 16
(continued)

Fig. 16
(continued)

cata	gcgc	ca	gac	ctt	gg	ttt	act	ttt	ctt	gt	act	ttt	gaatt	aca	agg	ttt	26500							
tgt	aaat	ttt	ggaaa	at	ttt	gtt	gtt	ttt	aaat	act	gt	gt	ttt	gct			26550							
ttt	aaat	aca	ac	att	c	t	c	g	at	at	ttt	gaga	att	gt	gt	ttt	26600							
AAC	CTA	ACAG	T	T	C	T	A	A	G	G	C	T	G	A	T	GGG	TC	CAG						
E	P	N	S	F	L	K	K	A	D	I	F	I	N	G	S	Q		26650						
TTA	GGAGA	AGA	GAT	TTT	TAT	TCA	ATT	GCAT	AA	ACT	TCT	AAG	AA	AGT	TCC	CAC	CTT	26700						
L	G	E	D	F	I	Q	L	H	K	L	L	R	K	S	T	F		26750						
CAAAA	ATG	CAA	AA	CT	TAT	GGT	CT	GAT	GT	TGG	TAG	GCT	CGA	AGA	AA	AGA								
K	N	A	K	L	Y	G	P	D	V	G	Q	P	R	R	K									
CGG	CTA	AGA	TG	TGA	AGA	Ggt	tag	ga	act	ag	agg	at	cg	aa	t	act	tt	26800						
T	A	K	M	L	K	S																		
ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	26850						
ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	26900						
ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	26950						
ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	27000						
cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	27050						
cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	27100						
cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	27150						
cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	cc	27200						
tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	27250						
tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	27300						
tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	27350						
tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	27400						
at	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	27450						
F	L	K	A	G	G	E	V	I	D	S	V													
ACAT	GGC	CAT	CA	Gta	agt	at	gt	tc	tc	t	at	t	act	tag	gaa	agta	agg	27500						
T	W	H	H																					
ctag	ctt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	27550						
aatt	tg	act	g	cgt	t	caa	ata	a	gaa	aa	aca	aa	at	gt	t	ct	ct	27600						
act	cc	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	27650						
tagt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	27700						
ccat	gg	aa	ac	at	at	a	aa	tt	aa	tt	aa	tt	aa	tt	aa	tt	aa	27750						
acc	cagg	gg	at	gt	ac	tt	aa	cc	aa	tt	gt	ttt	tt	ca	aa	ga	ca	27800						
gatt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	27850						
tctt	gg	tt	tt	tt	cc	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	27900						
tt	gg	tt	tt	tt	cc	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	27950						
tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	28000						
Y																								
TATT	TTG	AAT	GG	AC	GG	ACT	G	C	T	ACC	AG	GG	AA	GAT	TTT	CTAA	ACC	CTG	AT	GT	28050			
Y	L	N	G	R	T	A	T	R	E	D	F	I	N	P	D	V								
ATT	GG	AC	AT	TTT	T	TT	CAT	T	GT	G	AAA	AG	TTT	CC	AG	gt	aat	ag	tct		28100			
L	D	I	F	I	S	S	V	Q	K	V	F	Q												
ttt	aa	ttt	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	28150					
ttt	ct	ta	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	28200					
aa	ca	ac	at	g	ac	tt	at	cc	ac	tt	gt	tt	tt	cc	ct	ag	t	tt	28250					
tt	ac	cc	cc	at	gt	ac	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	28300					
taga	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	ttt	aa	28350					
tttcc	cc	ac	cc	ct	ac	tt	ca	tt	aa	aa	aa	aa	aa	tt	cc	aa	ga	cc	tt	aa	28400			
act	at	at	ga	ac	at	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	28450					
aa	gt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	28500					
gttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	ttt	28550					
V	V	V	V	V	V	V	V	V	V	V	V	V	V	V	V	V	V	28600						
GCACC	AG	GC	CT	GG	CA	AG	GA	AG	GT	CT	GG	TT	AG	G	AA	CA	AG	CT	TC	GC	CAT	AT	28650	
S	T	R	P	G	K	K	V	W	L	G	E	T	S	S	A	Y								
GGAG	GG	CC	G	GG	CC	CC	TT	G	C	T	AC	CC	TT	TT	G	C	AG	CT	GG	CT	TT	AT	Tgt	
G	G	G	G	A	P	L	L	S	D	T	F	A	A	G	F	M								
agt	ga	ag	cg	cg	ct	gg	cc	tt	ag	gg	tt	ca	gt	tg	ca	g	tc	tt	cc	at	ct	ct	28700	
tct	t	tt	t	tt	gt	ca	aa	tt	ag	ct	cc	cc	ag	cc	aa	tt	ag	cc	tt	cc	at	tt	28750	
tca	gt	tg	gt	ca	gg	cc	aa	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	tt	28800	
act	aa	ag	tt	tg	tt	gg	gg	ac	at	tt	ta	ca	ag	tg	tt	cc	aa	tt	aa	tt	act	ata	28850	
agg	at	gt	tt	tt	ca	ga	aa	tt	tt	gg	cc	tt	ta	tt	at	tt	gg	cc	cata	aa	at	at	28900	

Fig. 16
(continued)

Fig. 16
(continued)

Fig. 16
(continued)

tttgtcttgggtgatcatgtgtttgttatgtgtgtgtctaaaattt	34900
tggcttgagctttgtttgaattcttggatgaaacaataaccaagaatac	34950
ttaaactctgatcattctgacagatatcccacaggctatggctttt	35000
gaattgtgtcccccagtgataaaaaagcagcaagcacgatactgtctcag	35050
attcatggtggtcacatgtgaggtaaaaaaaaaaaaagatgaatccca	35100
ttaaatgcccccaggataacagtgtatcttttaggataactatgg	35150
cttgccactgggttattaaataaggacataagtaaagatctatgg	35200
ctcttctcccaaccaccacaactagGATTATTGGCTATCTCTGTGTT	35250
D Y W L S L L F	
CAAGAAATTGGTGGGCACCAAGGTGTTAATGGCACAGCGTGCAGGGTCAA	35300
K K L V G T K V L M A S V Q G S	
AGAGAAGGAAGCTCGAGTATACTTCATTGCACAAACACTGACAAgtaa	35350
K R R K L R V Y L H C T N T D N	
gtatgaaacacacccttaccaatcatcaagttttagtggtaagcctgt	35400
aactttactcaaacaccctgttgcattgtctatacattgcataagtata	35450
ggcagtgtcaatttagttaagtttatacaacgatttatTTTatTTat	35500
tttagaagaaaaatgctactttttgtgtgtgtttttgagacggggc	35550
ctcgctcgtccccaggctggagtgcaagtggtcaatctcagctcactgc	35600
aacctccgcctccgggtcaagtgtattctgaagaggagaacaataata	35650
acaacaatattttttcaaaagttgtaccgcagtctggagttgagaa	35700
gacatcgagatTTTgtccctataactttgttttaggttagcaaaaaat	35750
gttcttaaatctcaggataattttctatagtaggtttcaatctatcc	35800
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atttccgatttttttttttttttttttttttttttttttttttttttttttt	35900
gaagttagcgagggaaatgttttttttttttttttttttttttttttttt	35950
tagatTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	36000
ccccctgttttttttttttttttttttttttttttttttttttttttttttt	36050
gccttttttttttttttttttttttttttttttttttttttttttttttttt	36100
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ttggtagcatgaacggcaacatTTTTTTTTTTTTTTTTTTTTTTTTTT	36200
cactagcggctaaaacgatcataaaaagaaggatactaagagggcccact	36250
gtcattatggatcctaatacttaggatgcattatggattgtcattatgga	36300
tactaatacttaggatcacatTTTTTTTTTTTTTTTTTTTTTTTTTTTT	36350
agatacatatttttatttttttttttttttttttttttttttttttttttttt	36400
P R Y	
AAGAAGGAGATTTAACTCTGTATGCCATAAACCTCCATAATGTCACCAAG	36450
K E G D L T L Y A I N L H N . V T K	
TACTTGCCTTACCCCTATCCTTTCTAAACAGTGGATAAAATACCT	36500
Y L R L P Y F S N K Q V D K Y L	
TCTAAAGACCTTGGGACCTCATGGATTACTTCACAAgtaaatTTTCC	36550
L R P L G P H G L L S K	
ttgttcatccaaaacttcaataaaatttttttttttttttttttttttttt	36600
agtttggacaggaggcaaaagacaaagtcaactatatacaaggcttaataa	36650
ttcttaatattcaggaaatttttttttttttttttttttttttttttttttt	36700
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atcttagagaataaggTTTGTGTTTTTTTTTTTTTTTTTTTTTTTTTT	36800
actcaacgcatttttttttttttttttttttttttttttttttttttttttt	36850
acattcactaaagcaaaatatacTTTTTTTTTTTTTTTTTTTTTTTT	36900
tgggttgtataaaatataccatgtgagatcagtgtgtgttttttttttt	36950
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agtttacaggcTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	37250
gaagctgttttttttttttttttttttttttttttttttttttttttttttt	37300
ggcaatTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	37350
tgggtgttttttttttttttttttttttttttttttttttttttttttttt	37400
attgttttttttttttttttttttttttttttttttttttttttttttttttt	37450
ctgtcaccctgggggggggggggggggggggggggggggggggggggggg	37500
taaaaaaaaatTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	37550

Fig. 16
(continued)

Fig. 16
(continued)

Fig. 16
(continued)

ctttgtcagcaataatatgtgagaggacagattgttagatatgatagtat	43450
aaaaaaatggtaatgacaattcagaggcgaggagattctgtaaaactaaa	43500
attactataaatgaaattgttgcataagaggataaatttagaaaacac	43550
ccaatacctataactgtctgttaatgctgcctttctcaccttcctt	43600
ccttggttcagttggaaagctttggctgcaagtaacagaaaactccatat	43650
tcaaatgcttaagcaataaggaaatgtatattcccacataactagacgt	43700
tcaaacaggccaggctccagcacttcagtgctcaccaggatctgggtt	43750
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tctggtagcatgtatggctgttagctgtttcatggcccccttcaaaccat	43850
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tgaataactcttttcagagcttcacagcaaaccttcctcatgtctc	43950
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tggagaggggtgttgtcagttcacaaactgaacactgcagttctgcgtt	44100
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ttcaaatgttatgcctgttatggatatagtatctttaaaattttatTTT	44200
aatagctttaggggtacacactttgttacaggggtgaattgtgttagt	44250
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ttccattcaatttattcaatttaagtagatattttgttaaggagctaaagctg	44650
aaaattaaatttttagatcttcaataactcttaattttatatgttaagtgg	44700
tttttatattttcacatttgaataaagtagattttataaccttgatatt	44750
gtatgactattcttttagtaatgtaaagcctacagactcctacatttggaa	44800
accacttagtgtgtttcaccccttgttatactatcaggatcctcga	44898

**Fig. 16
(continued)**

human	MLLRSKPALP PPLMLLLLGP LGPLSPGALP RPAQAOQDVVD LDFFTQEPLH
mouse	~~~~~ ML RLLLLWLWGP LGALAQGAPA GTAPTDVVVD LEFYTKRPLR
rat	~~~~~ ~ LLLLWLWGR LRALTQGTPA GTAPTKDGVVD LEFYTKRLFQ

human	LVSPSFLSVT IDANLATDPR FLILLGSPKL RTLARGLSPA YLRFGGTKTD
mouse	SVSPSFLSIT IDASLATDPR FLTFLGSPPRL RALARGLSPA YLRFGGTKTD
rat	SVSPSFLSIT IDASLATDPR FLTFLSSPPRL RALSRGGLSPA YLRFGGTKTD

human	FLIEDPKKES TFEERSYWQS QVNQDICKYQ SIPPDVEEKL RLEWPYQEQL
mouse	FLIEDPDKEP TSEERSYWKS QVNHDICRSE PVSAAVLRKL QVEWPFOELL
rat	FLIEDPNNEP TSEERSYWQS QDNNDICGSD RVSADVL~~~

human	LLREHYQKKE KNSTYSRSSV DVLYTFANCS GLDLIFGLNA LLRTADLQWN
mouse	LLREQYQKEF KNSTYSRSSV DMLYSFAKCS GLDLIFGLNA LLRTPDLRWN
rat	~~~~~

human	SSNAQLLLDY CSSKGYNISW ELGNEPNSFL KKADIFINGS QLGEDYIQLH
mouse	SSNAQLLLDY CSSKGYNISW ELGNEPNSFW KKAHILIDGL QLGEDFVELH
rat	~~~~~

human	KLLRKSTFKN AKLYGPDVGQ PRRKTAKMLK SFLKAGGEVI DSVTWHHHYL
mouse	KLLQRSAFQN AKLYGPDIGQ PRGKTVKLLR SFLKAGGEVI DSLTWHHHYL
rat	~~~~~

human	NGRTATREDF LNPDVLDIFI SSVQKVFQVV ESTRPGKKVW LGETSSAYGG
mouse	NGRIATKEDF LSSDA LDTFI LSVQKILKVT KEITPGKKVW LGETSSAYGG
rat	~~~~~

human	GAPLLSDTFA AGFMWLDKLG LSARMGIEVV MRQVFFGAGN YHLVDENFDP
mouse	GAPLLSNTFA AGFMWLDKLG LSAQMGIEVV MRQVFFGAGN YHLVDENFEP
rat	~~~~~

human	LPDYWLSLLF KKLVGTKVLM ASVQGSKRRK LRVYLHCTNT DNPRYKEGDL
mouse	LPDYWLSLLF KKLVGPVRLL SRVKGPDRSK LRVYLHCTNV YHPRYQEGDL
rat	~~~~~

human	TLYAINLHNV TKYLRLPYPFF SNKQVDKYLL RPLGPHGLLS KSVQLNGLTL
mouse	TLYVINLHNV TKHLKVPPPL FRKPVDTYLL KPSGPDGLLS KSVQLNGQIL
rat	~~~~~

human	KMVDDQTLPP LMEKPLRPAGS SLGLPAFSYS FFVIRNAKVA ACI~
mouse	KMVDEQTLPA LTEKPLPAGS ALSLPAFSYG FFVIRNAKIA ACI~
rat	KMVDEQTDXPA LTEKPLPAGS SLSVPAFSYG FFVIRNAKIA ACI~

Fig. 17

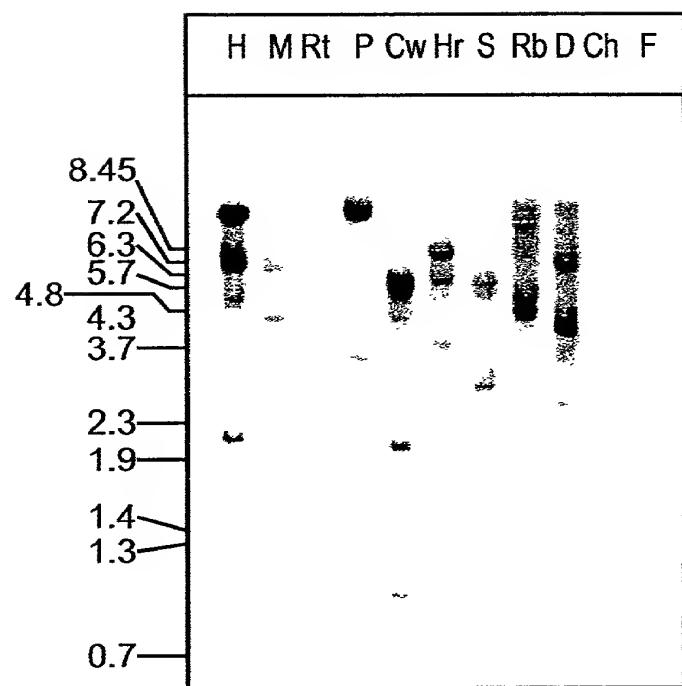


Fig. 18

Fig. 19